M. 2 ib 21/6

PAGE: 1

# RAW SEQUENCE LISTING PATENT APPLICATION US/09/234, 182A

DATE: 05/08/2000 TIME: 18:35:10

INPUT SET: \$35445.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

1 2		SEQUENCE LISTING ENTERED
3	(1)	General Information:
4 5 6 7 8	(i)	APPLICANT: Genentech, Inc., Hsei, Vanessa Koumenis, Iphigenia Leong, Steven R. Presta, Leonard G.
9 10 11		Shahrokh, Zahra Zapata, Gerardo A.
12 13 14	(11)	TITLE OF INVENTION: ANTIBODY FRAGMENT-POLYMER CONJUGATES AND HUMANIZED ANTI-IL-8 MONOCLONAL ANTIBODIES
15 16	(iii)	NUMBER OF SEQUENCES: 72
17 18 19 20 21 22 23 24	(iv)	CORRESPONDENCE ADDRESS: (A) ADDRESSEE: Genentech, Inc. (B) STREET: 1 DNA Way (C) CITY: South San Francisco (D) STATE: California (E) COUNTRY: USA (F) ZIP: 94080
25 26 27 28 29	(∨)	COMPUTER READABLE FORM:  (A) MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk  (B) COMPUTER: IBM PC compatible  (C) OPERATING SYSTEM: PC-DOS/MS-DOS  (D) SOFTWARE: WinPatin (Genentech)
31 32 33 34 35	(vi)	CURRENT APPLICATION DATA:  (A) APPLICATION NUMBER: 09/234182  (B) FILING DATE: 20-Jan-1999  (C) CLASSIFICATION:
36 37 38 39	(vii)	PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: 60/074330 (B) FILING DATE: 22-JAN-1998
40 41 42 43	(Vii)	PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: 60/094003 (B) FILING DATE: 24-JUL-1998
44 45 46	(Vii)	PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: 60/094013 (B) FILING DATE: 24-JUL-1998

#### RAW SEQUENCE LISTING PATENT APPLICATION US/09/234,182A

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```
47
48
     (vii) PRIOR APPLICATION DATA:
            (A) APPLICATION NUMBER: 60/075467
49
            (B) FILING DATE: 20-FEB-1998
50
51
52
    (viii) ATTORNEY/AGENT INFORMATION:
53
            (A) NAME: Love, Richard B.
            (B) REGISTRATION NUMBER: 34,659
54
            (C) REFERENCE/DOCKET NUMBER: P1085R4-1A
55
56
      (ix) TELECOMMUNICATION INFORMATION:
57
58
            (A) TELEPHONE: 650/225-5530
            (B) TELEFAX: 650/952-9881
59
60
    (2) INFORMATION FOR SEQ ID NO:1:
61
        (i) SEQUENCE CHARACTERISTICS:
62
            (A) LENGTH: 22 base pairs
63
64
            (B) TYPE: Nucleic Acid
65
            (C) STRANDEDNESS: Single
66
            (D) TOPOLOGY: Linear
67
      (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
68
69
70
71
     CAGTCCAACT GTTCAGGACG CC 22
72
    (2) INFORMATION FOR SEQ ID NO:2:
73
74
        (i) SEQUENCE CHARACTERISTICS:
75
76
            (A) LENGTH: 22 base pairs
77
            (B) TYPE: Nucleic Acid
78
            (C) STRANDEDNESS: Single
79
            (D) TOPOLOGY: Linear
80
81
      (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
82
8.3
     GTGCTGCTCA TGCTGTAGGT GC 22
84
85
    (2) INFORMATION FOR SEQ ID NO:3:
86
87
        (i) SEQUENCE CHARACTERISTICS:
88
89
            (A) LENGTH: 23 base pairs
90
            (B) TYPE: Nucleic Acid
91
            (C) STRANDEDNESS: Single
92
            (D) TOPOLOGY: Linear
93
94
      (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
95
96
97
     GAAGTTGATG TCTTGTGAGT GGC 23
98
99
    (2) INFORMATION FOR SEQ ID NO:4:
```

### RAW SEQUENCE LISTING PATENT APPLICATION US/09/234, 182A

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100	
101	(i) SEQUENCE CHARACTERISTICS:
102	(A) LENGTH: 24 base pairs
103	(B) TYPE: Nucleic Acid
104	(C) STRANDEDNESS: Single
105	(D) TOPOLOGY: Linear
106	
107	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
108	
109	
110	GCATCCTAGA GTCACCGAGG AGCC 24
111	
112	(2) INFORMATION FOR SEQ ID NO:5:
113	
114	(i) SEQUENCE CHARACTERISTICS:
115	(A) LENGTH: 22 base pairs
116	(B) TYPE: Nucleic Acid
117	(C) STRANDEDNESS: Single
118	(D) TOPOLOGY: Linear
119	
120	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
121	
122	
123	CACTGGCTCA GGGAAATAAC CC 22
124	
125	(2) INFORMATION FOR SEQ ID NO:6:
126	
127	(i) SEQUENCE CHARACTERISTICS:
128	(A) LENGTH: 22 base pairs
129	(B) TYPE: Nucleic Acid
130	(C) STRANDEDNESS: Single
131	(D) TOPOLOGY: Linear
132	
133	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
134	
135	
136	GGAGAGCTGG GAAGGTGTGC AC 22
137	
138	(2) INFORMATION FOR SEQ ID NO:7:
139	A CONTRACT ON PAGE TOTAL
140	(i) SEQUENCE CHARACTERISTICS:
141	(A) LENGTH: 35 base pairs
142	(B) TYPE: Nucleic Acid
143	(C) STRANDEDNESS: Single
144	(D) TOPOLOGY: Linear
145	(wi) GEOMENGE DECORPORANT GEO ID NO.7.
146	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
147	
148 149	ACAAACGCGT ACGCTGACAT CGTCATGACC CAGTC 35
150	ACAMACGEGI ACGETGACAT EGICATGACE CAGTE 35
151	(2) INFORMATION FOR SEQ ID NO:8:
151	(2) INCOMMITON FOR SEQ ID NO.0:

## RAW SEQUENCE LISTING PATENT APPLICATION US/09/234,182A

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INPUT SET: S35445.raw

153 (i) SEQUENCE CHARACTERIST 154 (A) LENGTH: 35 base p	
154 (A) LENGTH: 35 base p	airs
155 (B) TYPE: Nucleic Aci	d
156 (C) STRANDEDNESS: Sin	gle
157 (D) TOPOLOGY: Linear	-
158	
159 (xi) SEQUENCE DESCRIPTION:	SEO ID NO:8:
160	
161	
162 ACAAACGCGT ACGCTGATAT TGTCA	TGACT CAGTC 35
163	
164 (2) INFORMATION FOR SEQ ID N	0:9:
165	
166 (i) SEQUENCE CHARACTERIST	ICS:
167 (A) LENGTH: 35 base p	
168 (B) TYPE: Nucleic Aci	
169 (C) STRANDEDNESS: Sin	
170 (D) TOPOLOGY: Linear	320
171	
172 (xi) SEQUENCE DESCRIPTION:	SEO ID NO:9:
173	DIQ ID NO.
174	
175 ACAAACGCGT ACGCTGACAT CGTCA	TGACA CAGTC 35
176	
177 (2) INFORMATION FOR SEQ ID N	0:10:
178	
179 (i) SEQUENCE CHARACTERIST	TCS:
180 (A) LENGTH: 37 base p	
181 (B) TYPE: Nucleic Aci	
182 (C) STRANDEDNESS: Sin	
183 (D) TOPOLOGY: Linear	540
184	
185 (xi) SEQUENCE DESCRIPTION:	SEO ID NO:10:
186	DDQ 1D NOVIO
187	
188 GCTCTTCGAA TGGTGGGAAG ATGGA	TACAG TTGGTGC 37
189	inono ilicoloco o,
190 (2) INFORMATION FOR SEQ ID N	0:11:
191	· · · · · · · · · · · · · · · · · · ·
192 (i) SEQUENCE CHARACTERIST	TCS:
193 (A) LENGTH: 39 base p	
194 (B) TYPE: Nucleic Aci	
195 (C) STRANDEDNESS: Sin	
196 (D) TOPOLOGY: Linear	9-0
197	
198 (xi) SEQUENCE DESCRIPTION:	SEO ID NO:11:
199	DDQ ID NOTII.
200	
201 CGATGGGCCC GGATAGACCG ATGGG	GCTGT TGTTTTGGC 39
202	55.51 101111000 57
203 (2) INFORMATION FOR SEQ ID N	0:12:
203 (2) INFORMATION FOR SEQ ID N	V.12.
205 (i) SEQUENCE CHARACTERIST	TCS:
(1) Shyother ellatherest	100.

## RAW SEQUENCE LISTING PATENT APPLICATION US/09/234,182A

DATE: 05/08/2000 TIME: 18:35:11

INPUT SET: S35445.raw

206		(A) LENGTH: 39 base pairs
207		(B) TYPE: Nucleic Acid
208		(C) STRANDEDNESS: Single
209		(D) TOPOLOGY: Linear
		(b) for onodi: binear
210		
211	(X1)	SEQUENCE DESCRIPTION: SEQ ID NO:12:
212		
213		
214	CGATGO	GCCC GGATAGACTG ATGGGGCTGT CGTTTTGGC 39
215		
216	(2) TNE	FORMATION FOR SEQ ID NO:13:
	(2) 1111	ORMATION FOR SEQ ID NO.13.
217		
218	(1)	SEQUENCE CHARACTERISTICS:
219		(A) LENGTH: 39 base pairs
220		(B) TYPE: Nucleic Acid
221		(C) STRANDEDNESS: Single
222		(D) TOPOLOGY: Linear
223		(b) for obodi. Einear
		GROUPINGS PROGRESSION GROUP TO NO. 12.
224	(XI)	SEQUENCE DESCRIPTION: SEQ ID NO:13:
225		
226		
227	CGATGO	GGCCC GGATAGACGG ATGGGGCTGT TGTTTTGGC 39
228		
229	(2) TNE	FORMATION FOR SEQ ID NO:14:
	(Z) INI	ORMATION FOR SEQ ID NO.14.
230		
231	(1)	SEQUENCE CHARACTERISTICS:
232		(A) LENGTH: 39 base pairs
233		(B) TYPE: Nucleic Acid
234		(C) STRANDEDNESS: Single
235		(D) TOPOLOGY: Linear
236		(5) 101020011 21001
237	/ ** i \	SEQUENCE DESCRIPTION: SEQ ID NO:14:
	(XI)	SEQUENCE DESCRIPTION: SEQ ID NO:14:
238		
239		
240	CGATGO	GCCC GGATAGACAG ATGGGGCTGT TGTTTTGGC 39
241		
242	(2) INE	FORMATION FOR SEQ ID NO:15:
243	( - ,	
244	7 1 1	SEQUENCE CHARACTERISTICS:
	(1)	
245		(A) LENGTH: 39 base pairs
246		(B) TYPE: Nucleic Acid
247		(C) STRANDEDNESS: Single
248		(D) TOPOLOGY: Linear
249		
250	(vi)	SEQUENCE DESCRIPTION: SEQ ID NO:15:
	( A I )	Phyonych phyckiliton, phy in Mo.11.
251		
252		
253	CGATGO	GGCCC GGATAGACTG ATGGGGCTGT TGTTTTGGC 39
254		
255	(2) INE	FORMATION FOR SEQ ID NO:16:
256	, ,	~
257	/ i \	SEQUENCE CHARACTERISTICS:
	( _ )	
258		(A) LENGTH: 369 base pairs

### **SEQUENCE VERIFICATION REPORT** PATENT APPLICATION *US/09/234,182A*

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INPUT SET: S35445.raw

Line Error Original Text

Wrong application Serial Number (A) APPLICATION NUMBER: 09/234182